# Increased mitophagy in the skeletal muscle of spinal and bulbar muscular atrophy patients

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## **Supplementary Table 1**

SBMA patient	Age at onset (years)	Age at biopsy (years)	CAG repeats number	ALS patie		Age at onset (years)	Age at biopsy (years)	Nerurogenic atrophy	Sex	Age at onset (years)	Age at biopsy (years)	Control	Sex	Age at biopsy (years)
1	50	52	43	1	М	23	23	1	М	23	23	1	М	52
2	31	80	41	2	м	46	48	2	м	46	48	2	м	42
3	60	54	43	3	м	41	42	3	М	41	42	3	м	55
4	50	59	46	4	М	45	46	4	м	45	46	4	м	53
5	26	57	45	5	М	55	55	5	м	55	55	5	м	63
6	38	61	44	6	м	62	62	6	М	62	62	6	м	36
7	41	45	46	7	М	67	67	7	М	67	67	7	м	45
8	64	71	43	8	М	81	83	8	М	81	83	8	м	28
9	49	57	44	9	м	77	81	9	М	77	81	9	м	42
10	45	61	46	10	м	60	61	10	М	60	61	10	м	32
11	37	44	49	11	м	77	77	11	М	77	77	11	м	45
12	46	55	48	12	м	61	61	12	м	61	61	12	м	45
13	27	28	43	13	м	38	39	13	М	38	39	13	м	44
14	29	39	46	14	м	38	40	14	М	38	40	14	м	79
15	45	54	48	15	М	46	47					15	м	38
16	49	55	44	16	м	49	50					16	м	38
17	43	64	41	17	м	53	54					17	м	19
18	48	51	44	18	м	55	56					18	м	47
19	34	37	49	19	М	56	57					19	F	27
				20	М	55	57					20	F	38
				21	м	58	59					21	F	39
				22	М	63	64					22	F	44
				23	м	64	66					23	F	50
				24	М	67	67							
				25	М	67	68							
				26	М	73	73							
				27	F	30	33							
				28	F	45	48							
				29	F	51	51							
				30	F	50	55							
				31	F	65	65							
				32	F	64	66							

 $Table \ S1. \ List \ of \ muscle \ biopsy \ specimens \ collected \ from \ patients \ affected \ with \ SBMA, ALS, \ and \ other neurogenic \ diseases.$ 

## **Supplementary Table 2**

	Forward primers	Reverse primers	Probe
APP	5'-TTTTTGTGTGCTCTCCCAGGTCT-3'	5'-TGGTCACTGGTTGGTTGGC-3'	5'-CCCTGAACTGCAGATCACCAATGTGGTAG-3
AR	5'-TTGTCCACCGTGTGTCTTCTTCTGC-3'	5'-TGCACTTCCATCCTTGAGCTTGGC-3'	
COII	5'-CGTCTGAACTATCCTGCCCG-3'	5'-TGGTAAGGGAGGGATCGTTG-3'	5'-CGCCCTCCCATCCCTACGCATC-3'
COX4	5'-CATGTGGCAGAAGCACTATGTGT-3'	5'-GCCACCCACTCTTTGTCAAAG-3'	
CRLS1	5'-CCCAGTTCTGGGCTATTTGA-3'	5'-TCTTTGATTGGCCCAGTTTC-3'	
ERRα	5'-TTCTCATCGCTGTCGCTGTCT-3'	5'-CAGCCGCCGCACTAGTTG-3'	
Gp78	5'-ACCTCCTGTCCAACATGCAG-3'	5'-GCAATCCGAGACCCATCGAA-3'	
MFN1	5'-TGTTTTGGTCGCAAACTCTG-3'	5'-CTGTCTGCGTACGTCTTCCA-3'	
MFN2	5'-ATGCATCCCCACTTAAGCAC-3'	5'-CCAGAGGGCAGAACTTTGTC-3'	
MnSOD	5'-CTTCAGCCTGCACTGCCGTTCAAT-3'	5'-CTGAAGGTAGTAAGCGTGCTCCC-3'	
MUL1	5'-GCTGTTATAGAAGGAGCTGTGC-3'	5'-GCACTGTGTTGGTCCTCTGA-3'	
NRF1	5'-GGTGCAGCACCTTTGGAGAA-3'	5'-CCAGAGCAGACTCCAGGTCTTC-3'	
PGC-1α	5'-TCAGTCCTCACTGGTGGACA-3'	5'-TGCTTCGTCGTCAAAAACAG-3'	
PGC-1β	5'-CTGCTGGCCCAGATACACTGA-3'	5'-ATCCATGGCTTCATACTTGCT-3'	
RPLPO	5'-GTGATGTGCAGCTGATCAAGACT-3'	5'-GATGACCAGCCCAAAGGAGA-3'	
TFAM	5'-GAACAACTACCCATATTTAAAGCTCA-3'	5'-GAATCAGGAAGTTCCCTCCA-3'	

Table S2. List of primers used for real-time PCR analysis of gene expression.

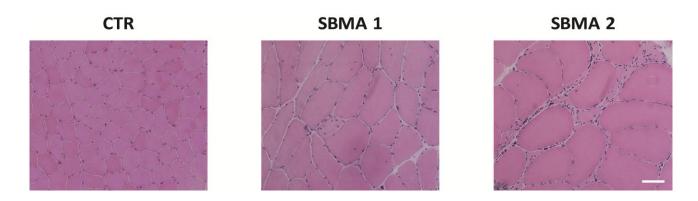


Figure S1. Atrophic and hypertrophic fibers in SBMA muscle.

H&E staining revealed the presence of small atrophic fibers together with enlarged, hypertrophic fibers in the quadriceps muscle of SBMA patients compared to age-matched control specimens. Shown are representative images from 1 control subject and 2 SBMA patients. Scale bar,  $80 \mu m$ .

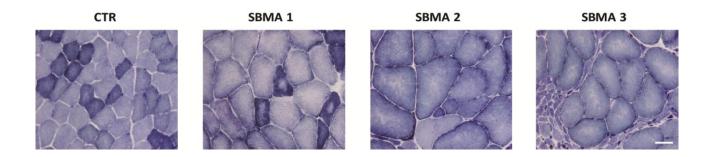


Figure S2. Both oxidative and glycolytic fibers are atrophic and hypertrophic in SBMA muscle.

NADH staining revealed the presence of small atrophic fibers together with hypertrophic glycolytic and oxidative fibers in the quadriceps muscle of SBMA patients compared to age-matched control specimens. Shown are representative images from 1 control subject and 3 SBMA patients. Scale bar,  $80 \mu m$ .

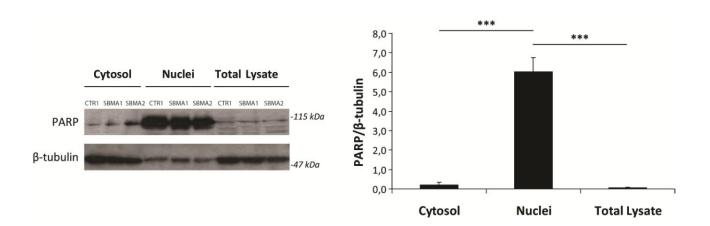


Figure S3. Purity of nuclear extracts.

Western blotting analysis of PARP (nuclear marker) and  $\beta$ -tubulin (cytosolic marker) in cytosolic fraction (cytosol), nuclear fraction (nuclei), and total lysates obtained from SBMA patient and control muscle samples. Nuclear enrichment was evaluated as the ratio between PARP and  $\beta$ -tubulin levels. Graph, mean  $\pm$  SEM, n = 2 SBMA patients and 1 control (CTR) subject. Significance by Student t test: \*\*\*p < 0.001.

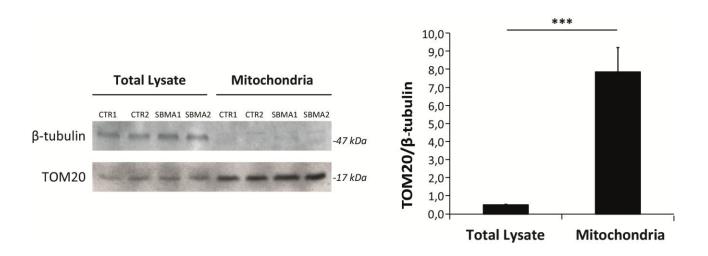


Figure S4. Purity of mitochondrial extract.

Western blotting analysis of TOM20 (mitochondrial marker) and  $\beta$ -tubulin (cytosolic marker) in muscle total lysates and isolated mitochondria derived from SBMA patient and control samples. Mitochondrial purification was evaluated as the ratio between TOM20 and  $\beta$ -tubulin levels. Graph, mean  $\pm$  SEM, n = 2 SBMA patients and 2 control (CTR) subjects. Significance by Student t test: \*\*\*p < 0.001.

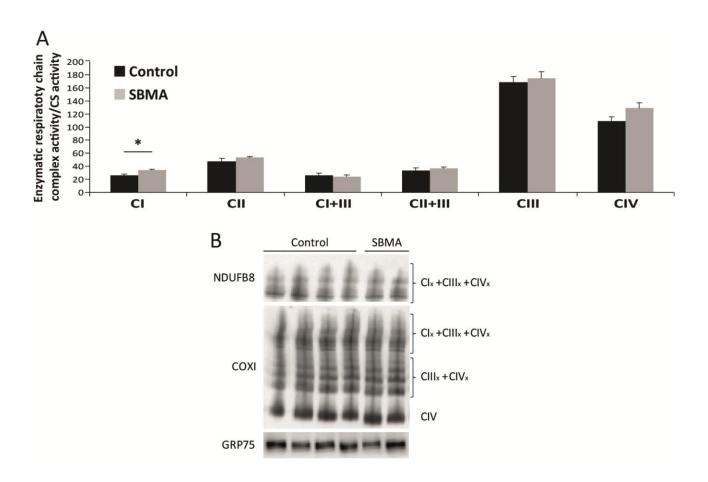


Figure S5. Normal OXPHOS activity upon normalization to CS activity and normal supercomplex expression and assembly in SBMA muscle.

- A) Enzymatic activity of respiratory chain complexes I–IV normalized to the CS activity. Graph, mean  $\pm$  SEM, n = 6 SBMA patients and 4 control subjects. Significance by Student *t* test: \*p < 0.05.
- B) Representative Blue Native PAGE (BN-PAGE) analysis of mitochondrial supercomplexes in isolated muscle mitochondria from 4 control subjects and 4 SBMA patients. Mitochondrial supercomplexes were recognized with antibodies against a subunit of Complex I (NDUFB8) and Complex IV (COXI). GRP75, a marker of mitochondrial matrix, was used as loading control.

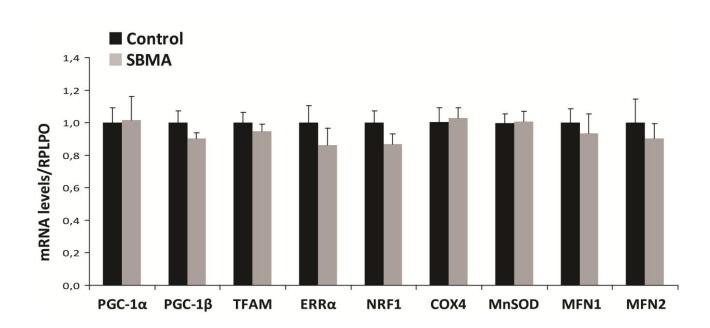


Figure S6. Normal expression of genes involved in mitochondrial biogenesis and function in muscles from SBMA patients. RT-PCR analysis of the transcript levels of genes involved in mitochondrial biogenesis and function, including peroxisome proliferator-activated receptor-gamma coactivator 1 alpha and 1 beta (PGC-1 $\alpha$  and PGC-1 $\beta$ ), mitochondrial transcription factor A (TFAM), estrogen-related receptor alpha (ERR $\alpha$ ), nuclear respiratory factor 1 (NRF1), cytochrome c oxidase 4 (COX4), manganese-dependent superoxide dismutase (MnSOD) and mitofusin 1 and 2 (MFN1 and MNF2), normalized to large ribosomal protein (RPLPO). Graph, mean  $\pm$  SEM, n = 14 SBMA patients and 18 control subjects.

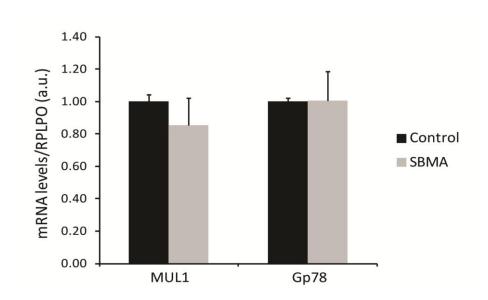


Figure S7. The expression of the E3 ubiquitin ligases, MUL1 and Gp78, is not altered in the muscle of SBMA patients. RT-PCR analysis of the transcript levels of genes involved in mitophagy, including MUL1 and Gp78. Graph, mean  $\pm$  SEM, n = 5 SBMA patients and 3 control subjects.